

## Amendment to the Specification.

Please replace the first full paragraph on page 6 (lines 5-16) with the following:

- -The invention provides a crystal comprising a polypeptide selected from (a) a modified Hdm2 protein, characterized by structural coordinates comprising a root mean square deviation of conserved residue backbone atoms of less than about 1.5 Å (e.g., less than about 1.0 Å, less than about 0.5 Å, or less than about 0.1 Å) when superimposed on backbone atoms described by structural coordinates of Table 3 or (b) a modified Hdm2 protein characterized by structural coordinates comprising a root mean square deviation of conserved residue backbone atoms of less than about 1.5 Å (e.g., less than about 1.0 Å, less than about 0.5 Å, or less than about 0.1 Å) when superimposed on backbone atoms described by structural coordinates of Table 4. In certain embodiments, the crystal may be complexed with a compound that binds to modified Hdm2 (e.g., SCH549128, Ac-<sup>6</sup>ClWAC<sub>3c</sub>E or Ac-<sup>6</sup>BrWAC<sub>3c</sub>E)- -

Please replace the last partial paragraph on page 45 (lines 28-30) with the following:

- -A 13 ml aliquot of the 3.6 mg/ml pool was concentrated to a 1.25 mM concentration (in 3ml), on an AMICON 5000 mwco Ultrafree membrane. A 3 ml aliquot of 8.5 mM SCH549128 Ac-<sup>6</sup>ClWAC<sub>3c</sub>E (MW= 535) in Buffer A was added to the- -